

09/981,876

203

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Val Pro Ser Pro Phe Gly Cys Met Ile Phe Phe Phe Phe Lys Asn Pro
 20 25 30
 Trp Lys Gln Arg Leu Leu Gln Gly Trp Leu Gly Ala Arg Pro Ile His
 35 40 45
 Leu Leu Gly Tyr Leu Pro Leu Ser Leu Leu Trp Cys Pro Phe Pro Leu
 50 55 60
 Pro Cys Ala Arg Cys Ser Val Val Tyr Ile Ser Ser Pro Arg His Gly
 65 70 75 80
 Ala His Ala Pro Arg Asp Met Ile Leu Ser Leu Val Leu Ala His Gly
 85 90 95
 Ala Leu Tyr Lys Glu Leu Gly Gly Arg Gly Arg Lys Trp Glu Pro Ser
 100 105 110

Xaa

<210> SEQ ID NO 200
 <211> LENGTH: 123
 <212> TYPE: PRT
 <213> ORGANISM: Homo sapiens

<400> SEQUENCE: 200

Met Ala Cys Arg Cys Leu Ser Phe Leu Leu Met Gly Thr Phe Leu Ser
 1 5 10 15
 Val Ser Gln Thr Val Leu Ala Gln Leu Asp Ala Leu Leu Val Phe Pro
 20 25 30
 Gly Gln Val Ala Gln Leu Ser Cys Thr Leu Ser Pro Gln His Val Thr
 35 40 45
 Ile Arg Asp Tyr Gly Val Ser Trp Tyr Gln Gln Arg Ala Gly Ser Ala
 50 55 60
 Pro Arg Tyr Leu Leu Tyr Tyr Arg Ser Glu Glu Asp His His Arg Pro
 65 70 75 80
 Ala Asp Ile Pro Asp Arg Phe Ser Ala Ala Lys Asp Glu Ala His Asn
 85 90 95
 Ala Cys Val Leu Thr Ile Ser Pro Val Gln Pro Glu Asp Asp Ala Asp
 100 105 110
 Tyr Tyr Cys Ser Val Gly Tyr Gly Phe Ser Pro
 115 120

<210> SEQ ID NO 201
 <211> LENGTH: 315
 <212> TYPE: PRT
 <213> ORGANISM: Homo sapiens
 <220> FEATURE:
 <221> NAME/KEY: SITE
 <222> LOCATION: (9)
 <223> OTHER INFORMATION: Xaa equals any of the naturally occurring L-
 amino acids
 <221> NAME/KEY: SITE
 <222> LOCATION: (311)
 <223> OTHER INFORMATION: Xaa equals any of the naturally occurring L-
 amino acids
 <221> NAME/KEY: SITE
 <222> LOCATION: (315)
 <223> OTHER INFORMATION: Xaa equals stop translation

<400> SEQUENCE: 201

Met Ala Gly Gly Arg Cys Gly Pro Xaa Leu Thr Ala Leu Leu Ala Ala
 1 5 10 15

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Trp Ile Ala Ala Val Ala Ala Thr Ala Gly Pro Glu Glu Ala Ala Leu
      20          25          30

Pro Pro Glu Gln Ser Arg Val Gln Pro Met Thr Ala Ser Asn Trp Thr
      35          40          45

Leu Val Met Glu Gly Glu Trp Met Leu Lys Phe Tyr Ala Pro Trp Cys
      50          55          60

Pro Ser Cys Gln Gln Thr Asp Ser Glu Trp Glu Ala Phe Ala Lys Asn
      65          70          75          80

Gly Glu Ile Leu Gln Ile Ser Val Gly Lys Val Asp Val Ile Gln Glu
      85          90          95

Pro Gly Leu Ser Gly Arg Phe Phe Val Thr Thr Leu Pro Ala Phe Phe
      100          105          110

His Ala Lys Asp Gly Ile Phe Arg Arg Tyr Arg Gly Pro Gly Ile Phe
      115          120          125

Glu Asp Leu Gln Asn Tyr Ile Leu Glu Lys Lys Trp Gln Ser Val Glu
      130          135          140

Pro Leu Thr Gly Trp Lys Ser Pro Ala Ser Leu Thr Met Ser Gly Met
      145          150          155          160

Ala Gly Leu Phe Ser Ile Ser Gly Lys Ile Trp His Leu His Asn Tyr
      165          170          175

Phe Thr Val Thr Leu Gly Ile Pro Ala Trp Cys Ser Tyr Val Phe Phe
      180          185          190

Val Ile Ala Thr Leu Val Phe Gly Leu Phe Met Gly Leu Val Leu Val
      195          200          205

Val Ile Ser Glu Cys Phe Tyr Val Pro Leu Pro Arg His Leu Ser Glu
      210          215          220

Arg Ser Glu Gln Asn Arg Arg Ser Glu Glu Ala His Arg Ala Glu Gln
      225          230          235          240

Leu Gln Asp Ala Glu Glu Glu Lys Asp Asp Ser Asn Glu Glu Glu Asn
      245          250          255

Lys Asp Ser Leu Val Asp Asp Glu Glu Glu Lys Glu Asp Leu Gly Asp
      260          265          270

Glu Asp Glu Ala Glu Glu Glu Glu Glu Asp Asn Leu Ala Ala Gly
      275          280          285

Val Asp Glu Glu Arg Ser Glu Ala Asn Asp Gln Gly Pro Pro Gly Glu
      290          295          300

Asp Gly Val Thr Arg Glu Xaa Ser Arg Ala Xaa
      305          310          315

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<110> SEQ ID NO 202
<211> LENGTH: 236
<212> TYPE: PRT
<213> ORGANISM: Homo sapiens
<220> FEATURE:
<221> NAME/KEY: SITE
<222> LOCATION: (236)
<223> OTHER INFORMATION: Xaa equals stop translation

<400> SEQUENCE: 202

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Met Gly Thr Ala Asp Ser Asp Glu Met Ala Pro Glu Ala Pro Gln His
  1          5          10          15

Thr His Ile Asp Val His Ile His Gln Glu Ser Ala Leu Ala Lys Leu
  20          25          30

Leu Leu Thr Cys Cys Ser Ala Leu Arg Pro Arg Ala Thr Gln Ala Arg

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